



ENTERED

1600

RAW SEQUENCE LISTING

DATE: 11/06/2002

PATENT APPLICATION: US/09/509,482A

TIME: 14:47:45

Input Set : D:\SEQLIST.txt

Output Set: N:\CRF4\11062002\I509482A.raw

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4 <110> APPLICANT: Crofts, Linda Anne
5      Hancock, Manuela S.
6      Morrison, Nigel A.
7      Eisman, John A.
9 <120> TITLE OF INVENTION: Isoforms of the Human Vitamin D Receptor
12 <130> FILE REFERENCE: RICE-014
14 <140> CURRENT APPLICATION NUMBER: 09/509,482A
15 <141> CURRENT FILING DATE: 2000-09-15
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18 <151> PRIOR FILING DATE: 1998-09-29
20 <150> PRIOR APPLICATION NUMBER: P09500
21 <151> PRIOR FILING DATE: 1997-09-29
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30 <213> ORGANISM: Homo sapiens
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47 gtgaaggctg caaaggcttc ttcaggcgaa gcatgaagcg gaaggcacta ttcacctgcc      360
48 ccttcaacgg ggactgccgc atcaccaagg acaaccgacg ccactgccag gcctgccggc      420
49 tcaaacgctg tgtggacatc ggcatgatga aggagtcatc totgacagat gaggaagtgc      480
50 agagggaagcg ggagatgata ctgaagcgga aggaggagga ggccttgaag gacagtctgc      540
51 ggcccaagct gctgtaggag cagcagcgca tcattgccat actgctggag gccaccata      600
52 agacctacga cccacacct tccgacttct gccagttccg gcctccagtt cgtgtgaatg      660
53 atggtggagg gagccatcct tccaggccca actccagaca cactcccagc ttctctgggg      720
54 actcctcctc ctctgtctca gatcactgta tcacctcttc agacatgatg gactcgtcca      780
55 gcttctccaa tctggatctg agtgaagaag attcagatga cccttctgtg accctagagc      840
56 tgtcccagct ctccatgctg cccacacctg ctgacctggg cagttacagc atccaaaagg      900
57 tcattggctt tgctaagatg ataccaggat tcagagacct cacctctgag gaccagatcg      960
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61 agaagctgaa cttgcatgag gaggagcatg tcctgtctcat ggccatctgc atcgtctccc 1200
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64 ccaagatgat ccagaagcta gccgacctgc gcagcctcaa tgaggagcac tccaagcagt 1380
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76 cttccctgcc tgaccctgga gactttgacc ggaacgtgcc ccgatctgt ggggtgtgtg 180
77 gagaccgagc cactggcttt cacttcaatg ctatgacctg tgaaggctgc aaaggcttct 240
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81 tgaagcggaa ggaggaggag gccttgaagg acagtctgcg gcccaagctg tctgaggagc 480
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95 ccgacctgcg cagcctcaat gaggagcact ccaagcagta ccgctgcctc tccttccagc 1320
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101 <212> TYPE: DNA

102 <213> ORGANISM: Homo sapiens

104 <400> SEQUENCE: 4

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106 ggctgtcgat ggtgtcaga actgtcggag tggaggggat ggaggcaatg gcggccagca 120
107 cttccctgcc tgaccctgga gactttgacc ggaacgtgcc ccgatctgt ggggtgtgtg 180
108 gagaccgagc cactggcttt cacttcaatg ctatgacctg tgaaggctgc aaaggcttct 240
109 tcagggtgagc cccctccca ggctctcccc agtggaaagg gagggagaag aagcaagggtg 300
110 tttccatgaa gggagccctt gcatttttca catctccttc cttacaatgt ccatggaaca 360
111 tgcggcgctc acagccacag gagcaggagg gtcttggcga agcatgaagc ggaaggcact 420

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113 ggcctgccgg ctcaaacgct gtgtggacat cggcatgatg aaggagttca ttctgacaga 540
114 tgaggaagtg cagaggaagc gggagatgat cctgaagcgg aaggaggagg aggccttgaa 600
115 ggacagtctg cggcccaagc tgtctgagga gcagcagcgc atcattgcca tactgctgga 660
116 cgcccaccat aagacctacg accccaccta ctccgacttc tgccagttcc ggcctccagt 720
117 tcgtgtgaat gatggtggag ggagccatcc ttccaggccc aactccagac aactcccag 780
118 cttctctggg gactcctcct cctcctgctc agatcactgt atcacctctt cagacatgat 840
119 ggactcgtcc agcttctcca atctggatct gagtgaagaa gattcagatg acccttctgt 900
120 gaccctagag ctgtcccagc tctccatgct gccccacctg gctgacctgg tcagttacag 960
121 catccaaaag gtcattggct ttgctaagat gataccagga ttcagagacc tcacctctga 1020
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123 gtccttcacc atggacgaca tgtcctggac ctgtggcaac caagactaca agtaccgcgt 1140
124 cagtgcagtg accaaagccg gacacagcct ggagctgatt gagccccctca tcaagttcca 1200
125 ggtgggactg aagaagctga acttgcataa ggaggagcat gtccctgctca tggccatctg 1260
126 catcgtctcc ccagatcgtc ctgggggtgca ggacgcgcgc ctgattgagg ccatccagga 1320
127 ccgcctgtcc aacacactgc agacgtacat ccgctgccgc cccccgcccc cgggcagcca 1380
128 cctgctctat gccaaagatga tccagaagct agccgacctg cgcagcctca atgaggagca 1440
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169 gggagatgat cctgaagcgg aaggaggagg aggccttgaa ggacagtctg cggcccaagc 660
170 tgtctgagga gcagcagcgc atcattgcca tactgctgga cgcccaccat aagacctacg 720
171 accccaccta ctccgacttc tgccagttcc ggccctccagt tcgtgtgaat gatggtggag 780
172 ggagccatcc ttccaggccc aactccagac acactcccag cttctctggg gactcctcct 840
173 cctcctgctc agatcaactgt atcacctctt cagacatgat ggactcgtcc agcttctcca 900
174 atctggatct gagtgaagaa gattcagatg acccttctgt gaccctagag ctgtcccagc 960
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183 tccagaagct agccgacctg cgcagcctca atgaggagca ctccaagcag taccgctgcc 1500
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190 <213> ORGANISM: Homo sapiens

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195 gg 122

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198 <211> LENGTH: 477

199 <212> TYPE: PRT

200 <213> ORGANISM: Homo sapiens

202 <400> SEQUENCE: 9

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206 20 25 30
207 Pro His Arg Arg Ala Pro Leu Gly Ser Thr Tyr Leu Pro Pro Ala Pro
208 35 40 45
209 Ser Gly Met Glu Ala Met Ala Ala Ser Thr Ser Leu Pro Asp Pro Gly
210 50 55 60
211 Asp Phe Asp Arg Asn Val Pro Arg Ile Cys Gly Val Cys Gly Asp Arg
212 65 70 75 80
213 Ala Thr Gly Phe His Phe Asn Ala Met Thr Cys Glu Gly Cys Lys Gly
214 85 90 95
215 Phe Phe Arg Arg Ser Met Lys Arg Lys Ala Leu Phe Thr Cys Pro Phe
216 100 105 110
217 Asn Gly Asp Cys Arg Ile Thr Lys Asp Asn Arg Arg His Cys Gln Ala
218 115 120 125
219 Cys Arg Leu Lys Arg Cys Val Asp Ile Gly Met Met Lys Glu Phe Ile

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223 Lys Glu Glu Glu Ala Leu Lys Asp Ser Leu Arg Pro Lys Leu Ser Glu
224      165      170      175
225 Glu Gln Gln Arg Ile Ile Ala Ile Leu Leu Asp Ala His His Lys Thr
226      180      185      190
227 Tyr Asp Pro Thr Tyr Ser Asp Phe Cys Gln Phe Arg Pro Pro Val Arg
228      195      200      205
229 Val Asn Asp Gly Gly Gly Ser His Pro Ser Arg Pro Asn Ser Arg His
230      210      215      220
231 Thr Pro Ser Phe Ser Gly Asp Ser Ser Ser Ser Cys Ser Asp His Cys
232 225      230      235      240
233 Ile Thr Ser Ser Asp Met Met Asp Ser Ser Ser Phe Ser Asn Leu Asp
234      245      250      255
235 Leu Ser Glu Glu Asp Ser Asp Asp Pro Ser Val Thr Leu Glu Leu Ser
236      260      265      270
237 Gln Leu Ser Met Leu Pro His Leu Ala Asp Leu Val Ser Tyr Ser Ile
238      275      280      285
239 Gln Lys Val Ile Gly Phe Ala Lys Met Ile Pro Gly Phe Arg Asp Leu
240      290      295      300
241 Thr Ser Glu Asp Gln Ile Val Leu Leu Lys Ser Ser Ala Ile Glu Val
242 305      310      315      320
243 Ile Met Leu Arg Ser Asn Glu Ser Phe Thr Met Asp Asp Met Ser Trp
244      325      330      335
245 Thr Cys Gly Asn Gln Asp Tyr Lys Tyr Arg Val Ser Asp Val Thr Lys
246      340      345      350
247 Ala Gly His Ser Leu Glu Leu Ile Glu Pro Leu Ile Lys Phe Gln Val
248      355      360      365
249 Gly Leu Lys Lys Leu Asn Leu His Glu Glu Glu His Val Leu Leu Met
250      370      375      380
251 Ala Ile Cys Ile Val Ser Pro Asp Arg Pro Gly Val Gln Asp Ala Ala
252 385      390      395      400
253 Leu Ile Glu Ala Ile Gln Asp Arg Leu Ser Asn Thr Leu Gln Thr Tyr
254      405      410      415
255 Ile Arg Cys Arg His Pro Pro Pro Gly Ser His Leu Leu Tyr Ala Lys
256      420      425      430
257 Met Ile Gln Lys Leu Ala Asp Leu Arg Ser Leu Asn Glu Glu His Ser
258      435      440      445
259 Lys Gln Tyr Arg Cys Leu Ser Phe Gln Pro Glu Cys Ser Met Lys Leu
260      450      455      460
261 Thr Pro Leu Val Leu Glu Val Phe Gly Asn Glu Ile Ser
262 465      470      475
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267 <213> ORGANISM: Homo sapiens
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VERIFICATION SUMMARY

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